

# Introduction to High Performance Computing (HPC) – Session 1

## using the "Computational Shared Facility" (CSF)

Course materials / slides available from:

<https://ri.itservices.manchester.ac.uk/course/rcsf/>

Research Platforms, IT Services

[its-ri-team@manchester.ac.uk](mailto:its-ri-team@manchester.ac.uk)

<https://ri.itservices.manchester.ac.uk/csf3/>

Course materials at <https://ri.itservices.manchester.ac.uk/course/rcsf/>

## Feedback

- Your feedback is important to us!
- Please give feedback on this course
  - Quick form at <https://goo.gl/forms/zfZyTLw4DDaySnCF3>  
(choose "Introduction to HPC (Using CSF)")
  - Feedback is important to help us improve our courses
  - Records your attendance on the course

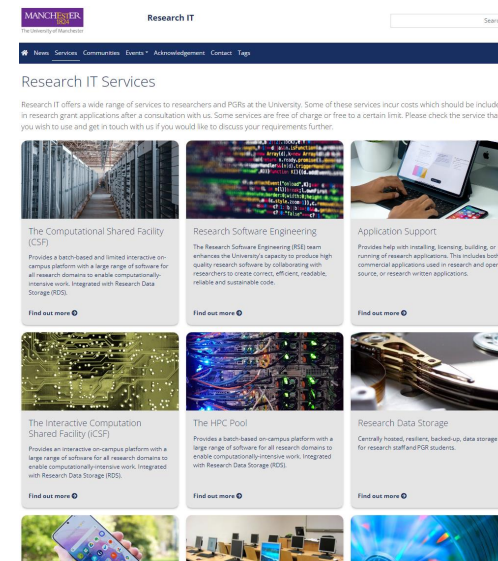
## Housekeeping

- Please let me know if you're leaving
  - 10am - 12:30pm (practical sessions 1, 2, & 3)
  - Lunch *approx* 12:30pm - 1:30pm
  - 1:30pm - 4pm (practical sessions 4 & 5)
- 1-to-1 help is available if needed during exercises.
- Power adapters are available for the purposes of charging laptops, please be considerate of other users and *be careful of any trailing leads*.
- Got a question at any point? PLEASE ASK!!

Course materials at <https://ri.itservices.manchester.ac.uk/course/rcsf/>

## Who we are - Research IT

<https://research-it.manchester.ac.uk/services/>



## What we'll cover today

- Brief intro to High Performance Computing (HPC) as motivation
- *Using* the University's HPC system - The "Computational Shared Facility" (CSF)
  - What the CSF is and what it can do for you
  - Logging in
  - Running work ("jobs") on the system
  - Different types of jobs (simple to advanced)
  - Using real applications
  - Doing the above in the practical sessions today

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Course materials at <https://ri.itservices.manchester.ac.uk/course/rcsf/>

## Who the course is for - everyone

- People new to HPC / research computing, or who just want to try the CSF
  - We'll introduce you to these topics and you'll try it out today
- Maybe your supervisor asked you to get a CSF account
  - We'll teach you how to use it
- Those who have used CSF already, but want to know more
  - Parallel jobs, job arrays, the batch system, ...
- *Using* the CSF is today's focus, so that you can use it effectively in your work
  - *not* theoretical aspects of HPC
    - but we'll explain some of the basics to help you make good use of the resources
  - *not* parallel software development or version control
    - But we'll show you how to run high-end parallel applications
  - *not* Linux installation / administration
    - but we'll cover the basic Linux commands needed to use the CSF
  - *not* the specifics of the software you plan on using
    - ask your PI/supervisor for help with that

Invest a little time now, get results much faster!

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<https://ri.itservices.manchester.ac.uk/course/rcsf/>

## Motivation: Why use HPC (and the CSF)?

- Some (most?) research computation not suitable for your desktop/laptop
  - Takes *too long* to run
  - Needs more memory
  - Uses *too much* disk/storage space
- Use advanced, centrally-managed, UoM hardware
- Eventually, use regional / national supercomputers

Do not let the size/capacity/power of your computer dictate the size and complexity of the models/simulations/systems/problems you are solving!

## WHY & WHAT ...

High Performance Computing : why use it & what is it

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# What is HPC?

- insideHPC.com
  - High Performance Computing most generally refers to the practice of **aggregating computing power** in a way that delivers **much higher performance** than one could get out of a typical desktop computer or workstation **in order to solve large problems** in science, engineering, or business.
- HPC systems are usually a **cluster of compute nodes** (with some extra items such as **login nodes**, storage, networking)
  - The CSF fits this description

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# A new way of working!



## Running on a desktop

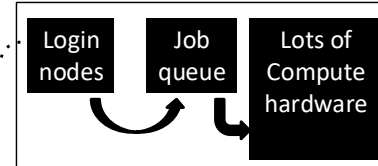
- You can fire up a GUI, run an app immediately. BUT:
- Got enough memory, cores, storage?
- Need to keep the PC to yourself (public cluster PC?)
- For several days?!
- Only one "job" (simulation, analysis) at a time?



## Logging in to the HPC system

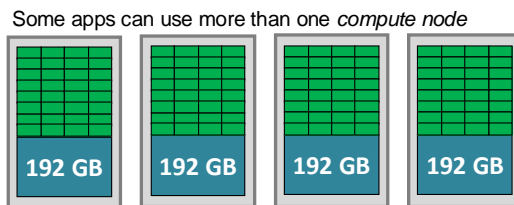
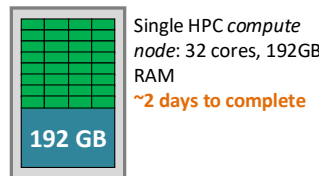
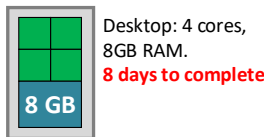
- Submit "**jobs**" to the queue
- Jobs wait until selected to run
- Jobs run on **high-end** hardware (lots of cores, memory, disk, GPU)
- Jobs run safely for days
- Many jobs can run at once
- Can log out any time (jobs still run)
- Log in to check on progress, get results

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## HPC Example: Finite Element Analysis

- Perform stress analysis on 3D mesh
  - The app splits the **input** into chunks
  - It performs calculations on chunks, **in parallel**
    - Faster and/or larger problem size

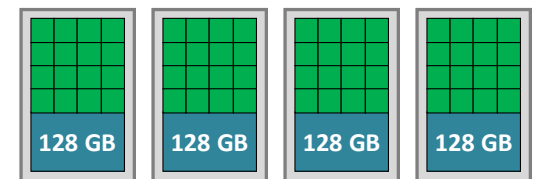
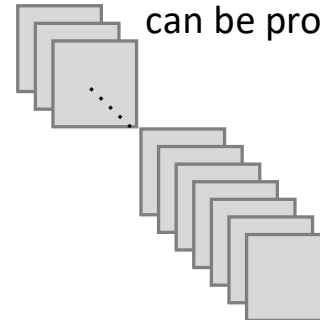
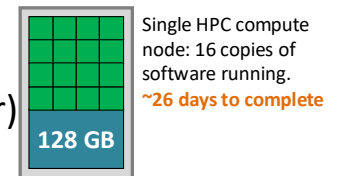
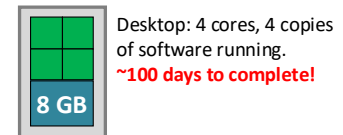
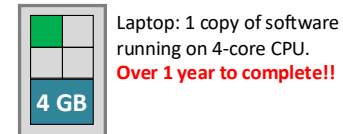


Multiple HPC compute nodes:  
128 cores, 768GB RAM  
~0.5 days to complete

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## "HTC" Example: Image Analysis

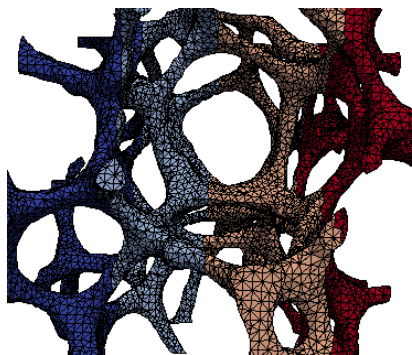
- High **Throughput** Computing
  - Not all s/w is "HPC" / parallel
  - But you might have **lots** of data
  - EG: Each image takes 1hr to process (and are independent - can be processed in any order)



Multiple HPC compute nodes: 64 copies of software running independently.  
~6 days to complete

Example: 10,000 image scans to be analysed by an image processing application. Each image takes 1 hour to process.

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Source: Professor Paul Mummery (MACE) using ParaFEM software <http://parafem.org.uk>

# What we'll be using today - the CSF

- Q: who has used the Computational Shared Facility (CSF) before?
- CSF3 current config:
  - A large Linux cluster system
  - 14,016 CPU cores (Intel “Xeon” or AMD “EPYC” CPUs)
  - 152 Nvidia GPUs (68 x v100, 72 x A100, 12 x L40s)
  - Got big datasets to process? Can run large-memory jobs
  - (we'll cover all of these details throughout the course)

BUT, you do not need to be running huge parallel jobs, or be a Linux / HPC expert, to use our systems and to benefit from the CSF

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# Who can use the CSF?

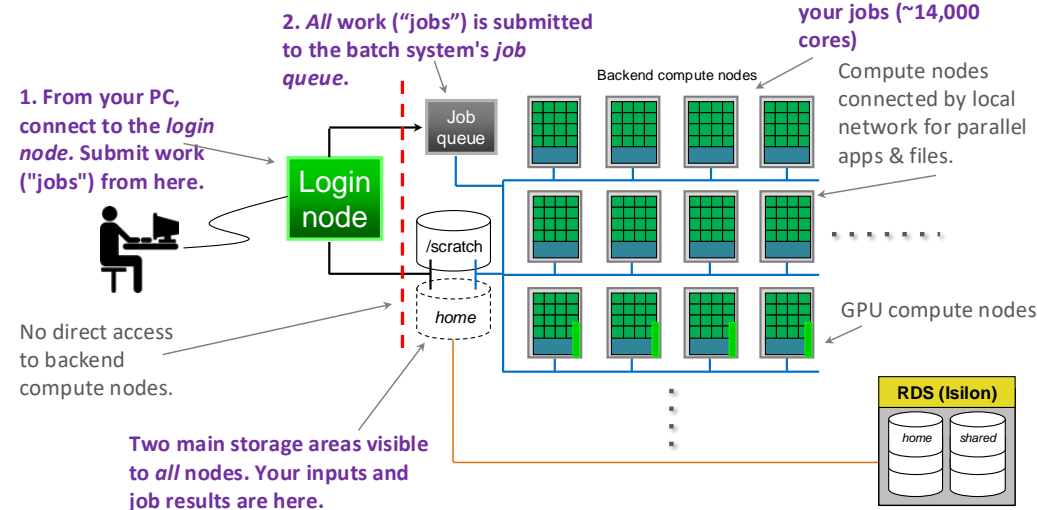
The following info is mainly for people who may want to "buy in" to the CSF. Your PI/supervisor or School may well have already done this! **If interested, ask us at the end of the course.**

- CSF uses a *shared funding model*
  - Researchers/academics/schools *contribute financially* to buy compute hardware
  - All h/w pooled so that all users can access the h/w
  - H/w *not* associated with individuals so it can always be in use as long as there are jobs to run!
- The time it takes to run all of your jobs depends on the size of the contribution with which you are associated
  - A research group that contributes more will be able to get more jobs done *sooner*
  - Managed automatically by the *batch system* – you just submit jobs!
- Some limited 'free at the point of use' access for non-contributors

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# What is the CSF? (more details)

- Computational Shared Facility
- A *batch* compute *cluster* to run your "jobs" (simulations, analysis,...)
- Here are the main components you'll learn about:



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## CSF: THE BASICS...

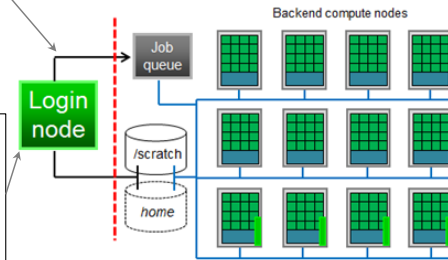
Hardware, OS, logging in, security, home filesystem, copying files, Linux, GUIs

## Login Nodes

- We really only see the CSF login node
  - Approximately 400 compute nodes
  - Far too many for *you* to find a free compute node to use
  - Instead: connect to the login node, let *job queue* find a node

### Key concept!

- Do:** Submit work ("jobs") to be run
  - No direct access to compute nodes
  - Submit jobs to the *job queue*.
  - The system will run your work on available *compute nodes* meeting your requirements.
- Do not:** run applications on the login node
  - Shared by all users, not much memory
  - For *lightweight* tasks (job submission, file transfer, ...)

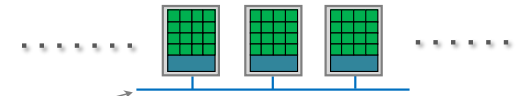


## Compute Nodes

- The CSF is a cluster of powerful "*compute nodes*"
  - Can think of the compute nodes as **very high-end** PCs where your simulations / data analysis / ML training ... will run
  - Different types of compute nodes available to suit your requirements (high mem, GPU, or a standard / default node)

- Multi-core CPUs (e.g., 16, 28, 32, 168 cores)
- Lots of RAM (e.g., 128GB, ..., 2TB, 4TB!)
- Network (possibly fast *InfiniBand* n/w)
- OS (Linux)
- Local disk (for temp files)
- Maybe a GPU

- Not all nodes have the same hardware.
- However, the CSF hides these details – let it choose the nodes your jobs use. You can optionally specify certain requirements.

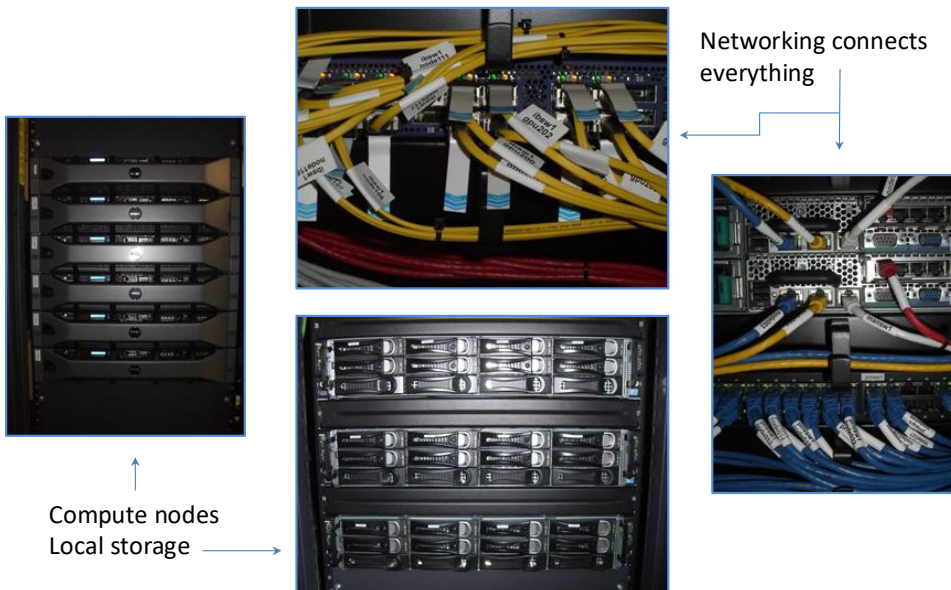


Local network connecting compute nodes. This allows:

- some applications to use more than one node (e.g., for big data / large simulations)
- all nodes to see all of your files - **so you don't have to copy files to the compute nodes**

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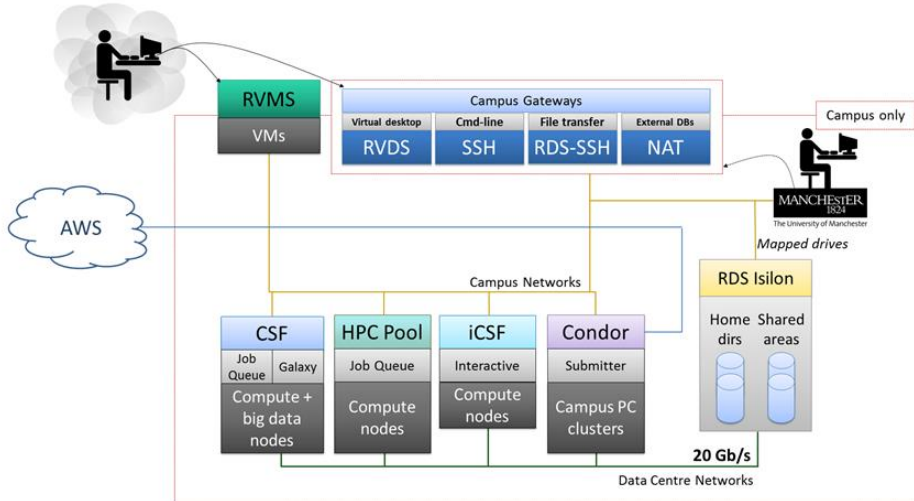
## Some pictures of the CSF



## Other Features of the CSF

- Variety of software applications (120+) & compilers
  - Gromacs, MATLAB, Abaqus, Bowtie2, Gaussian, Fluent, PyTorch + many more
- 1-core ("serial") jobs or *many*-core ("parallel") jobs
  - Do more work. Get it done sooner (see later!)**
- Lots of RAM: 128GB, 192GB, 512GB, 1.5TB, 2TB, 4TB(!!) compute nodes
- Lots of cores:
  - 16-core, 28-core or 32-core Intel compute nodes
  - 168-core AMD compute nodes (new Oct 2024!)
- Backed-up file storage (**no more USB disks!!!!!!!!!!**)
- Hardware failures don't stop your research
- Leave computational work running (for days)
- 14,000+ cores currently in the system
- Dedicated support team

# Part of the *Computationally Intensive Research Ecosystem*



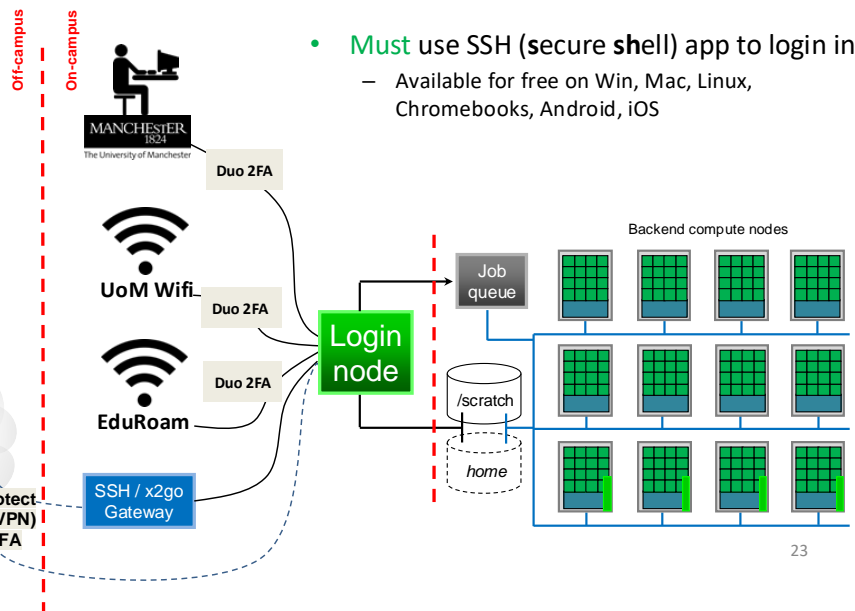
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## Logging in ...

Let's get access to the system

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## Where can I log in from?



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## Security (1)

- The CSF has a private (campus-only) IP address
  - Firewall also controls connections to and from the system
- When you are on-campus
  - Connect from any PC/laptop with a wired connection, or UoM WiFi, or EduRoam WiFi
  - Does not matter if using GlobalProtect or not, but you will *always* be asked to authenticate using your 2FA (DUO) device, e.g.

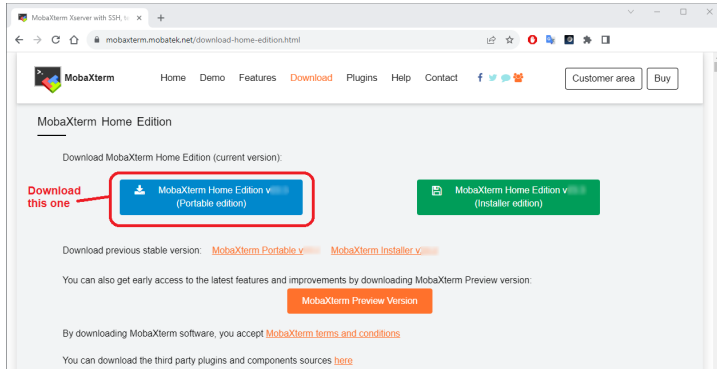
```

$ ssh -X myname@csf.itsservices.manchester.ac.uk
Password:
Duo two-factor login for myname@
Enter a passcode or select one of the following options:
  1. Duo Push to +XX XXXX XX6353
Passcode or option (1-1): 1
    
```

- When you are off-campus
  - First, sign-in to University GlobalProtect VPN + DUO 2FA
  - Then can login as normal to CSF (won't have to DUO 2FA again)
- Further documentation:
  - <https://ri.itsservices.manchester.ac.uk/csf3/getting-started/connecting/>

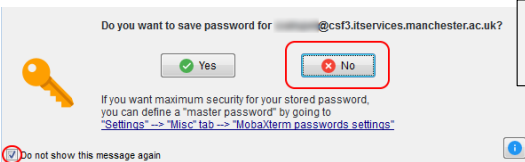
# Connect to CSF from Windows

- Access the CSF from a PC / laptop using an SSH (Secure Shell) app
  - Sometimes called a "terminal".
  - There's no web-site or other fancy GUI on the CSF – use the "command-line".
- Windows users** need to install a free *terminal* app called **MobaXterm**
- <https://mobaxterm.mobatek.net/download-home-edition.html>  
the **Home edition** (portable edition) does *not* require Administrator rights - just *extract* the small .zip file in your P-Drive or USB stick for example.



- Download using the blue box.
- Once downloaded, *right-click* on the .zip file and select:  
  
"Extract all ..."  
  
This will *unpack* the .zip file to a folder.

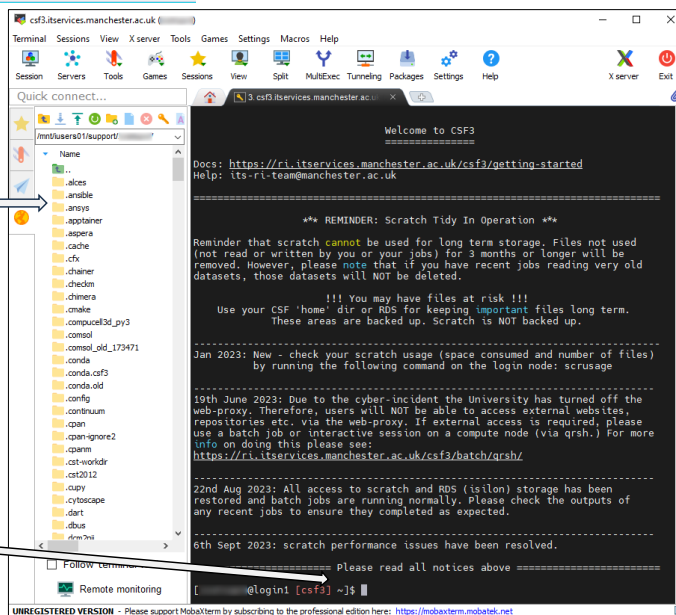
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If asked to save your password, we recommend you say "No", for security.

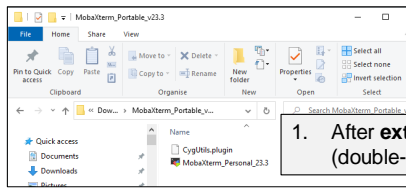
Drag-n-drop file browser for upload / download  
  
(new users won't have as many items in the list!)

We're on (one of) the CSF login nodes. Any commands you use will be typed "at the prompt", which shows your username and current directory (folder.)

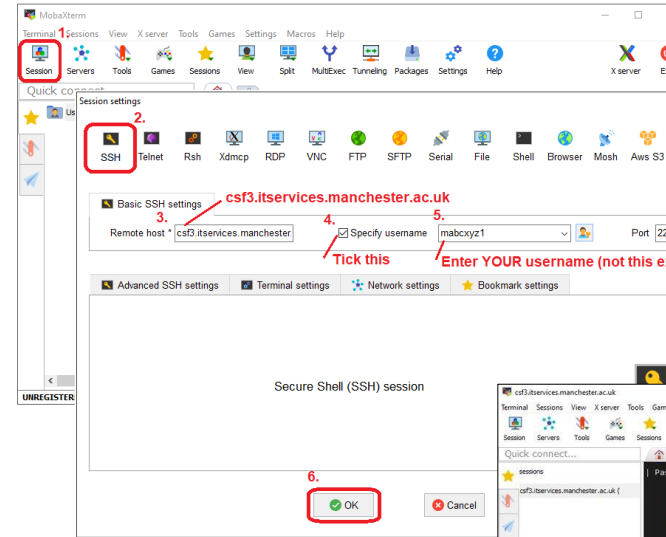


# MobaXterm "Session"

(username saved in the session setup)



- After **extracting** the .zip file, start MobaXterm\_Personal\_xy.z (double-click on the icon)



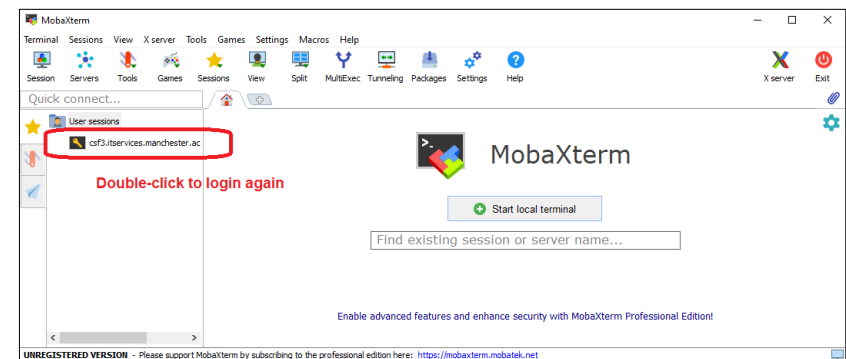
- 2 (1-6). Create a "Session" which saves the CSF's details along with *your* username.  
  
This is needed to make file drag-n-drop work (see later.)

3. This will then start to log you into the CSF – it will ask for your password. Type carefully!

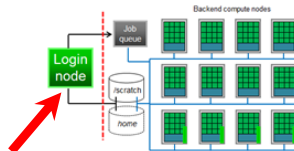
4. See slide about 2FA – you may be asked for DUO after your password

# Next time you want to login to CSF from Windows

- Just double-click the csf3 "session" in the list of "User sessions"
- The CSF details are saved in the "session"
- (this also makes the file browser work, for drag-n-drop file transfers.)



## Connect to CSF from Linux



- Access the CSF from a PC / laptop using an SSH (Secure Shell) app, eg a 'terminal'.
  - There's no web-site or other fancy GUI on the CSF - command-line for now.
- Linux users** - have a *terminal* application by default
  - Start a Terminal and type the following command:

```
ssh -X username@csf3.itservices.manchester.ac.uk
```

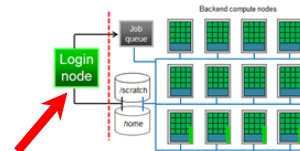
UPPERcase X

Central IT Services username.  
Answer 'Yes' to continue *if* asked.  
Enter central IT password when asked (same as for email)

- Finished using CSF? Log out with: `logout` or `exit`

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## Connect to CSF from a Mac



- Access the CSF from a PC / laptop using an SSH (Secure Shell) app, eg a 'terminal'.
  - There's no web-site or other fancy GUI on the CSF - command-line for now.
- Mac users** - have a *terminal* application by default
  - You will need to install X-Quartz first  
<https://www.xquartz.org/> (install then you should reboot your Mac)
  - Start a *Terminal* app and type the following command:

```
ssh -Y username@csf3.itservices.manchester.ac.uk
```

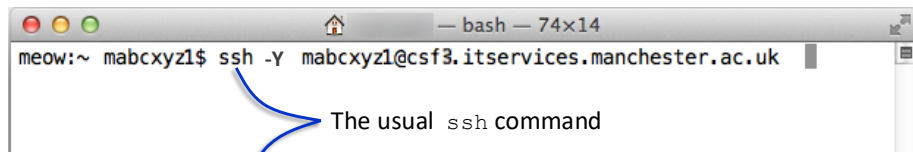
UPPERcase Y

Central IT Services username.  
Answer 'Yes' to continue *if* asked.  
Enter central IT password when asked (same as for email)

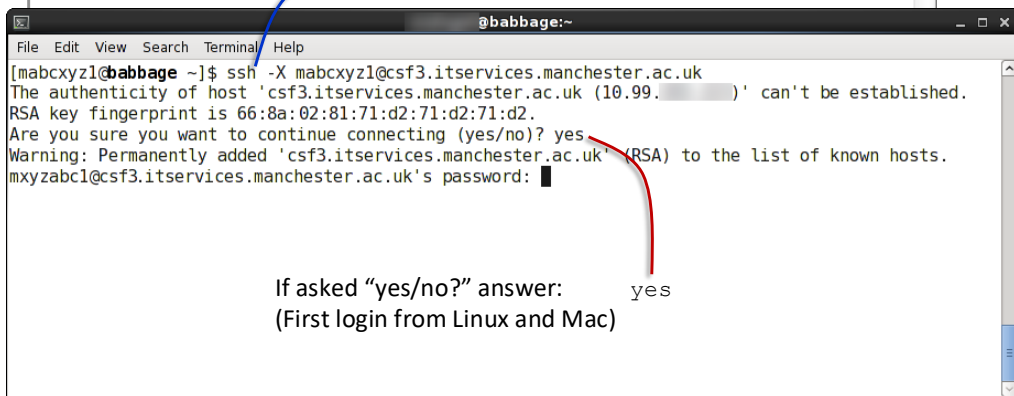
- Finished using CSF? Log out with: `logout` or `exit`

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## Linux / Mac Terminals



The usual `ssh` command



If asked "yes/no?" answer: yes  
(First login from Linux and Mac)

## DUO 2FA (when on-campus)

- When on-campus, after you enter your password, *all* login methods will then ask about DUO:

Duo two-factor login for *mabcxyz1*

Enter a passcode or select one of the following options:

1. Duo Push to +XX XXXX XX7890

Passcode or option (1-1): **1**  
Success. Logging you in...

(the message of the day is now displayed)

[mabcxyz1@login1 [csf3] ~]\$

Type 1 (and press Enter) in your ssh app to generate a DUO *push* to your device.

Then **accept** the push on your device.

You are now logged in.

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# What you see when you log in

- CSF uses CentOS Linux (c.f. Red Hat EL)
  - Command line – **requires the input of commands**, can be a little scary at first to new users
  - A welcome *message of the day* - announcements
  - The system awaits input/commands from you at a *prompt* (after you've logged in):

[username@login1 [csf3] ~]\$  
or [username@login2 [csf3] ~]\$

Type your  
commands  
at "the prompt"

- Learning Linux commands (more later):
  - <https://www.chm.bris.ac.uk/unix/>

# Security (2)

- It is **NOT** permitted to share your CSF account
- CSF uses your **IT password** – i.e. same as needed to access UoM email, Blackboard and so on ...
  - NEVER share it with ANYONE, including IT staff and your supervisor
  - Forgotten it? You can reset it via the IT Account Manager. Will affect all systems that require it.
    - <https://iam.manchester.ac.uk/>
- Reminder: Other general safety measures
  - Install a virus scanner  
<https://www.itservices.manchester.ac.uk/cybersecurity/advice/virusprotection/>
  - Be aware of phishing emails  
<https://www.itservices.manchester.ac.uk/cybersecurity/advice/phishing/>

## Practical Session 1 – Logging in

- Exercise 1 sheet (pdf) available at:  
<https://ri.itservices.manchester.ac.uk/course/rcsf/>
- Tip: During login, when you are prompted for your password, type carefully – you will not get a cursor that moves or display any \*\*\* as you press the keys. But it IS noticing what you type!
- Tip: Once logged in to the CSF, Linux does not always display something after you have entered a command – this is usually a good sign – your command worked, or there was nothing for it to do. If you've got something wrong, it will *usually* tell you via an error message.
- By the end of this practical session, everyone should have successfully logged in to the CSF!
- PLEASE ASK FOR HELP IF YOU RUN IN TO ANY PROBLEMS – WE ARE HERE TO HELP!

## PRACTICAL SESSION 1

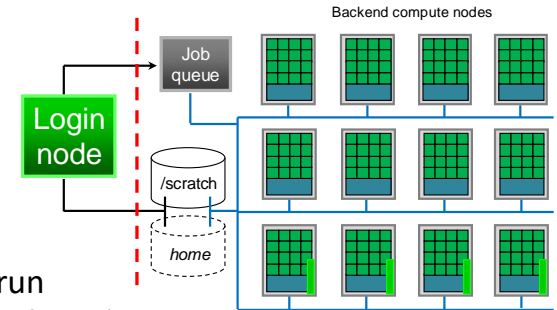
Logging in

## RUNNING JOBS

Doing real work on the CSF

## Jobs, Jobscripts and the Batch System

- We want to do computational work - “jobs”



- You decide:
  - Which program(s) to run
  - Which directory to run from (within *scratch* :-)
  - Which resources it needs (#cores, CPU type, memory)
- Write these requirements in a *jobscript*
- Submit your *jobscript* to the batch system (SGE)
- SGE decides exactly *when* and *where* the job runs

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## Reminder: The login nodes

- Do *not* run computational work here:
  - Not enough cores
  - Not enough memory
  - 100+ users connected, so running work causes serious problems
- You *can* do the following:
  - Transfer files on and off the CSF
  - Set up and submit your jobs (covered in next few slides)
  - Basic data processing/viewing
- Computational work running on the login node will be killed without warning!

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## Creating Jobscript files (1)

- You need to be able to create a small **text file** to describe your job
- Run `gedit` on the CSF login node - a simple text editor
  - Creates and saves the file *on the CSF*
  - `gedit` is similar to notepad (other Linux editors: nano, emacs, vi)
- Once logged in to the CSF type:  
`gedit &` '&' allow you to carry on using the command-line. Try it without to see.
  - Navigate to a file, or start typing, and then save

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## Can I Write Jobscripts on Windows?

- A warning about Windows text files (EG: in *notepad*)
  - There's an inconsistency over the (hidden) *end-of-line* characters in text files:
    - Windows: CR (carriage return) + LF (line feed)
    - Linux/Unix: LF (line feed)
  - The extra CR from Windows is a problem in jobscripts. Causes your job to **fail** immediately.
- Solutions
  - Use *gedit* on the CSF login node (writes Linux text files)
  - Or use *notepad*, upload then run `dos2unix myfile.txt`
    - Use only on jobscripts. **Do not come to rely on this – it is too easy to forget to do it – use gedit!**

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## A simple Jobscript – *Serial* (1 core)

myjob.txt

```
#!/bin/bash --login
#$ -cwd
#$ -N myjob
#$ -l resource
# Let's do work
date
hostname
sleep 120
date
```

Annotations:

- #!** on first line only (a special line)
- First line indicates we use the *bash* script language to write our jobscript.
- #\$** indicates a **batch system parameter** to specify our job requirements. We'll use various combinations of these.
- cwd** indicates we'll run from our current (working) directory. Input / output files will usually be found here.
- N (optional)**. Set the *jobname*. Otherwise, will use *name of your jobscript* as the job name.
- l (optional)** used to add extra resource requirements e.g. memory, time limits
- #** lines are just comments - anything on the line after it will be ignored.
- Actual Linux commands we run in our job. They will execute on a compute node.
- #\$ -l** course only works on the day of a course.

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## Submit Jobscript to Job Queue

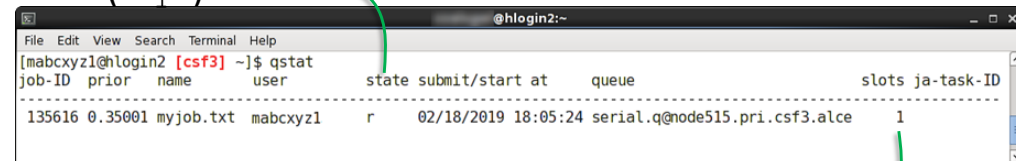
- Submit the jobscript from the login node with:  
`qsub jobscript` # EG: `qsub myjob.txt`
- You will be given a unique *JobID* (6/7-digit number)  
Your job 598052 ("*myjob.txt*") has been submitted
- You can then:
  - carry on with other work, submit other jobs that run other applications without disturbing previous jobs
  - log out of the CSF and your jobs will still run

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## Check status of your jobs

- To see your job(s) in the batch system, run:  
`qstat`

state is either queued (*qw*), running (*r*) or error (*Eqw*)



job-ID	prior	name	user	state	submit/start at	queue	slots	ja-task-ID
135616	0.35001	myjob.txt	mabcxyz1	r	02/18/2019 18:05:24	serial.q@node515.pri.csf3.alce	1	

slots is the number of cores (1 by default - a *serial* job)

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## Serial (1-core) Job Properties

- Our simple example job:
  - Serial (unless specified, only **1** core is used)
  - Standard memory (*no* **#\$** line asking for *more*)
    - We get: ~5GB RAM (per core - we're just using one core!)
  - Standard 7 days runtime (*no* **#\$** line asking for *shorter*)
- Default H/W: **standard serial jobs** will be placed on: **Intel nodes**
  - Remember, other nodes exist e.g., AMD, but you must add flags to your jobscript to get them
  - The system looks for a free core on a list of compute node that have been configured to run serial jobs.
- We *didn't* use any: **#\$ -l option**
  - EG: **#\$ -l mem2000** to land on higher memory node

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## Optional Serial Job Resources

<https://ri.itservices.manchester.ac.uk/csf3/batch/serial-jobs>

Jobscript flag(s)	Description
<b>#\$ -l short</b>	5GB/core Intel "haswell" CPU, <b>1 hour runtime limit</b> (for test/dev)
<b>#\$ -l mem512</b>	<b>32GB/core</b> Intel "haswell" CPU (usual 7 days runtime limit)
<b>#\$ -l mem512 -l short</b>	<b>32GB/core</b> Intel "haswell" CPU, <b>1 hour runtime limit</b> (for test/dev)
<b>#\$ -l amd -l short</b>	8GB/core <b>AMD EPYC "Genoa"</b> , <b>1 hour runtime limit</b> (for test/dev)
	Note: AMD nodes are mainly for parallel (multicore) jobs – see later. You <i>cannot</i> submit a serial (1-core), 7-day runtime job.
<b>#\$ -l mem1500</b>	<b>46GB/core</b> Intel "skylake" or "cascadelake" CPU (7-days runtime)
<b>#\$ -l mem2000</b>	<b>62GB/core</b> Intel "icelake" CPU (7-days runtime)

- **7-days runtime limit** on jobs, unless otherwise indicated in table.
  - "short" jobs have a **1-hour runtime**. They generally wait for less time in the queue.
- Our simple jobscript did *not* use any of the above. **Not needed in most cases.**
- If you limit a job by *node-type* or *memory* it may **wait longer in the queue**.
- You will see that the example jobscripts in the exercises have: **#\$ -l course**
  - **Only for use today** (we have reserved nodes on a teaching day.)
  - **Remove** if practicing after today (jobs will wait forever otherwise.)

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## So where did my results go?

- If `qstat` returns no output - means job has finished!
- Three possibilities:
  1. If app prints to screen: A text file called **jobname.oJobID**
    - *Jobname* is the name of your jobscript script (or `-N name` setting)
    - *JobID* is the number of your job
    - Previous example: `myjob.txt.o598052`
    - (Technically: "the std output stream is redirected to the file")
  2. An output file specific to your application
  3. Your job had a problem or failed: it may be reported in one of the above files or in **jobname.eJobID**
    - (Technically: "the std error stream is redirected to the file")
- Various options to view the file (they are plain text):
  - `cat filename`
  - `less filename` (allows you to page through with spacebar)
  - `gedit filename` (not recommended if it is large)

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## Why is my job still waiting?

- Your job will **wait** until there are cores available (meeting your jobscript's requirements).
- All jobs can run for up to 7 days (wallclock)
  - (**#\$ -l short** jobs up to 1 hour)
  - Jobs are starting and finishing all the time
- Initially frustrating (perhaps) but **advantages**:
  - You can log off, switch off your PC and your job will stay on the CSF. Log in later to check on job / collect the results.
  - You can submit many jobs.
    - They might not all run at once but SGE will decide...
  - More than one may run at the same time (make sure you have different files/folders for each job).

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## Many Users Sharing the CSF

- 100s of users running 1000s of jobs
- SGE gives each job a priority (number), which depends on
  - Size of research group's / school's CSF contribution
  - Amount of work already put through by that group and by you as an individual (this month)
- The time for your job to start depends on
  - Priority
  - Availability of requested resource (is CSF busy?)
- Jobs submitted *after* yours may start *before* yours!
- A few Jobs may **never** start
  - SGE tries to spot errors in jobscripts when you run qsub
  - Some may still get through then never run
- We try to ensure that if you submit some jobs, some of them will start within **24 hours**.
  - We make a check every morning of the waiting jobs

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## How busy is it?

- The system is usually **very** busy.
- However, jobs frequently finish, allowing waiting ones to start
- To see all the jobs for everyone

```
qstat -u "*"

```
- Note: all jobs shown as one list by qstat
  - It displays running and waiting jobs
  - Your job is not necessarily stuck behind all others above yours in the qstat output.
  - CSF is split into a few chunks – the very big jobs do not compete with the smaller jobs for cores
- **Do not try to guess when is a good time to submit your jobs.**
  - If you have work ready to run, **submit** it
  - If your jobs are not in the queue the scheduler cannot consider them
  - **You will waste time, not gain it, by not submitting**

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## What does Eqw mean? How do I delete a job?

- qstat reports your job as 'Eqw'
  - System tried to *start* it, but something went wrong
  - Usually no output file (job hasn't run) to indicate what happened, use:

```
qstat -j jobID | grep error

```
- Error can be cryptic. Most common causes:
  - Missing directory (cannot chdir to ...)
  - You created your jobscript on windows (exec error)
  - Unusual characters or **spaces** in file and directory names
  - No disk space on the filesystem - did you forget to use scratch?
- Detailed advice:  
<https://ri.itservices.manchester.ac.uk/csf3/batch/job-monitoring>
- To delete an Eqw job (it'll never run), or one you simply no longer want:

```
qdel jobID

```

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## PRACTICAL SESSION 2

Serial job

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## Practical Session 2 - Submitting jobs

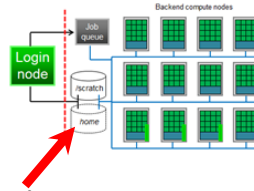
- Follow the handout 'Practical Session 2'
  - Use `qsub` to submit a simple serial job on the CSF
  - Use `qstat` to look at the queues
  - Use `qdel` to kill jobs
  - Use `qacct` to look at finished jobs
- Exercise sheet (pdf) available at:  
<https://ri.itservices.manchester.ac.uk/course/rcsf/>

## CSF STORAGE (FILESYSTEMS)

Where to store your files...

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### Storage – Home filesystem



- Upon login, automatically placed in your *home* directory (folder)  
`/mnt/iusers01/group01/username`
- Limited space, quota shared by everyone in the group
- Uses the Research Data Service (networked storage)
  - Large files can be slow-ish to read/write (implications for jobs)
- Which directory (folder) am I currently in?  
`pwd`
- How much space am I using? (Linux commands!)  
`du -sh dirname` # Can take a while
- How big is that file?  
`ls -lh filename` # Letter el not number l
- How much space is used/free overall?  
`df -h .` # The . is important!

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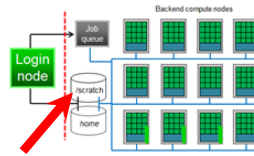
### Storage - Home filesystem

- *Home* is **backed up** and **mirrored** to another datacentre
  - Keep **important** files here (results, jobscripts, source code, ...)
  - Deleted a file by mistake? [its-ri-team@manchester.ac.uk](mailto:its-ri-team@manchester.ac.uk) can tell *you* how to retrieve it
- Only you can access your home directory
  - File permissions can be used to give others access
  - Contact [its-ri-team@manchester.ac.uk](mailto:its-ri-team@manchester.ac.uk) if you want advice on this as they can be complex
- **Do not** run jobs from your *home* area (see later)
  - Can generate a lot of files, some of them large
  - Using up all of the shared space will make your colleagues unhappy!
  - Consider compressing large (text) files with `gzip`

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## Filesystems - Scratch



- Filesystem local to CSF for:
  - Temporary files - can be huge
  - Running jobs from (it is faster!). **Recommended!**
- Shared by all CSF users, but we have 1.2PB
- Tidy up after each job finishes
- **Clean-up policy applies: files that have not been accessed for the past 3 months may be deleted automatically**
- **Not backed up!**
  - Move/copy important results to *home* area
  - Not considered safe for long term storage - hardware failure could cause data loss

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## Filesystems - Scratch

- Using scratch is easy: after log in, change to it:
 

```
cd ~/scratch
```

  - Uses a 'symlink' (short cut) in your home dir to `/scratch/username`
- Create a directory (now we're in scratch):
 

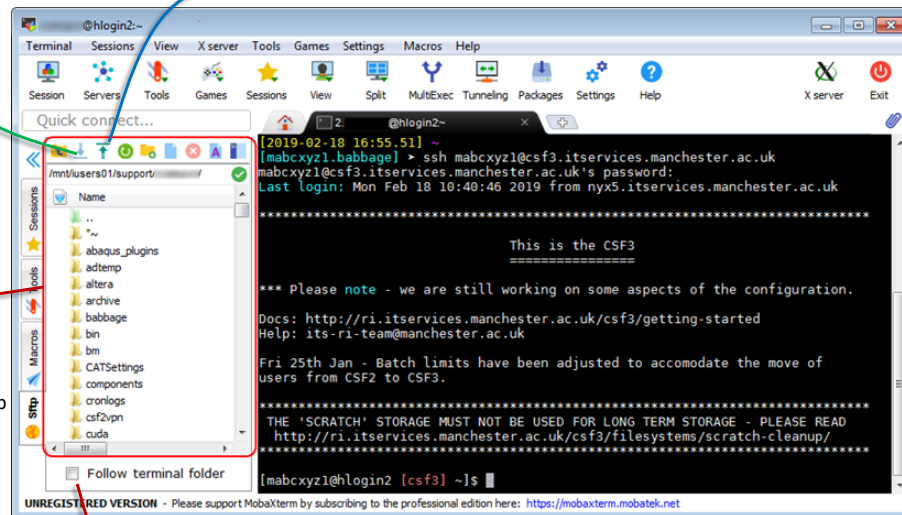
```
mkdir myjobdir
```
- Put all files relevant to your job in that directory and *run your jobs* from there - we'll try this out soon...
- **All compute nodes** see the same scratch area

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## File transfer with MobaXterm

2. First select files in the MobaXterm browser. Then the **Download** button opens a file-browser to select a destination folder on your PC.

3. The **Upload** button opens a file-browser to select files on your computer to **upload** to CSF (current directory).

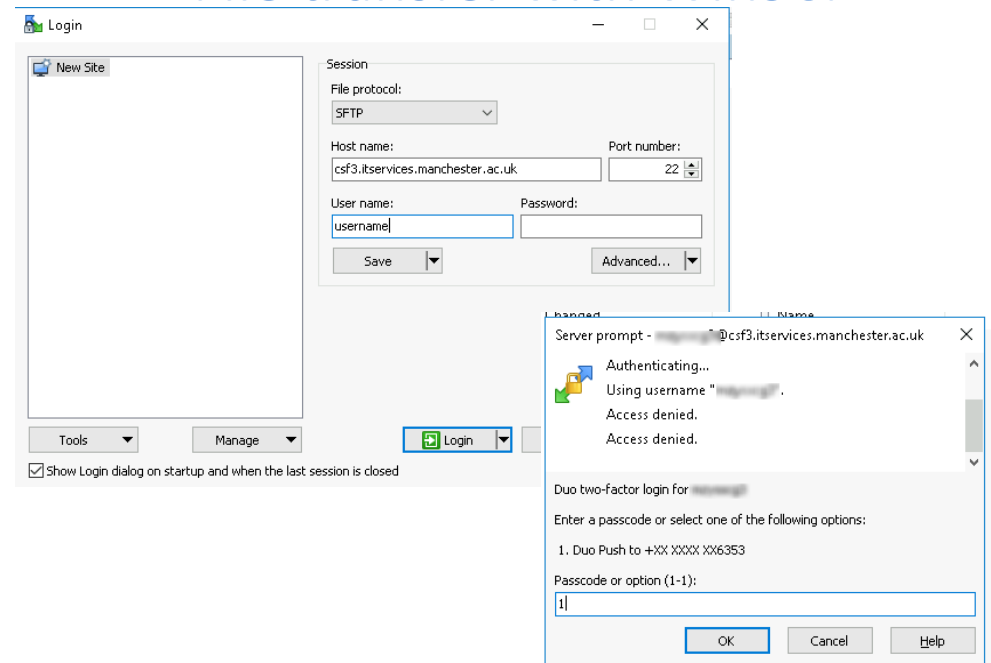


1. Drag-n-drop files from Windows Explorer or Desktop to/from here.

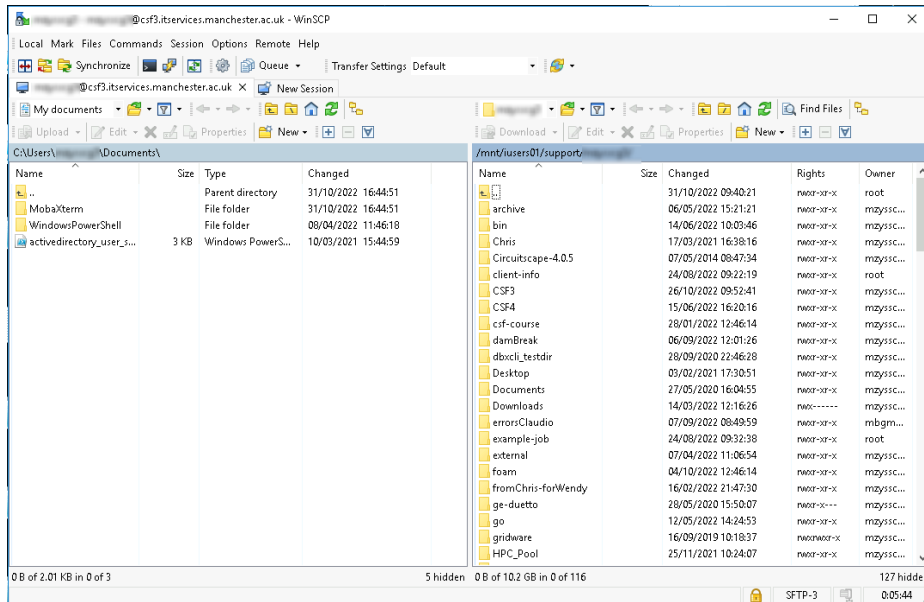
This can be flaky - do not use

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## File transfer with WinSCP



# File transfer with WinSCP



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# home file transfer with Linux / Mac (can also type in a local MobaXterm window)

- Transfer a file from your computer to your CSF home dir

`scp myfile.txt username@csf3.itservices.manchester.ac.uk:~/training/`

Exercise: Create a file on your PC named myfile.txt containing some text then transfer it to the CSF.

Destination directory on the CSF  
~ is shorthand meaning "your home directory"  
If no destination after the : then uses "your home directory"

- Transfer a file from your CSF home dir to your computer

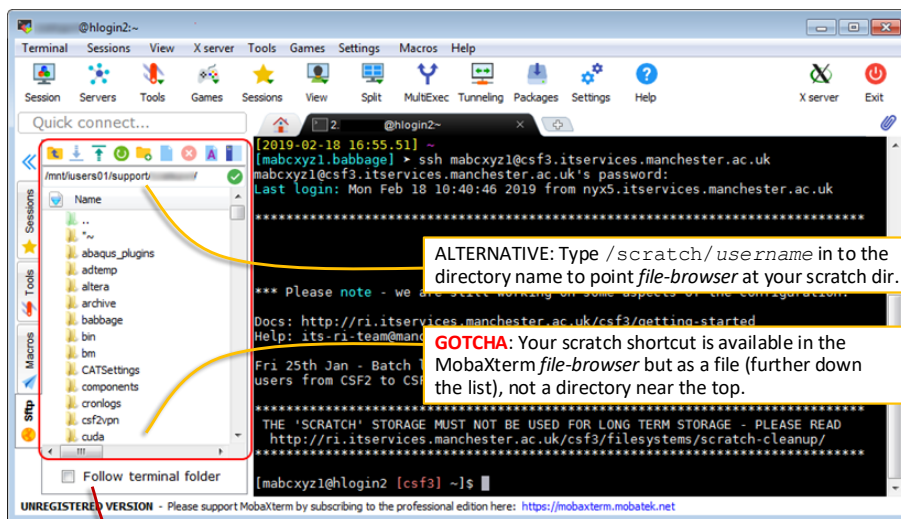
`scp username@csf3.itservices.manchester.ac.uk:results.out .`

The . is shorthand meaning "the current directory" on your computer

- Change directory & filenames...

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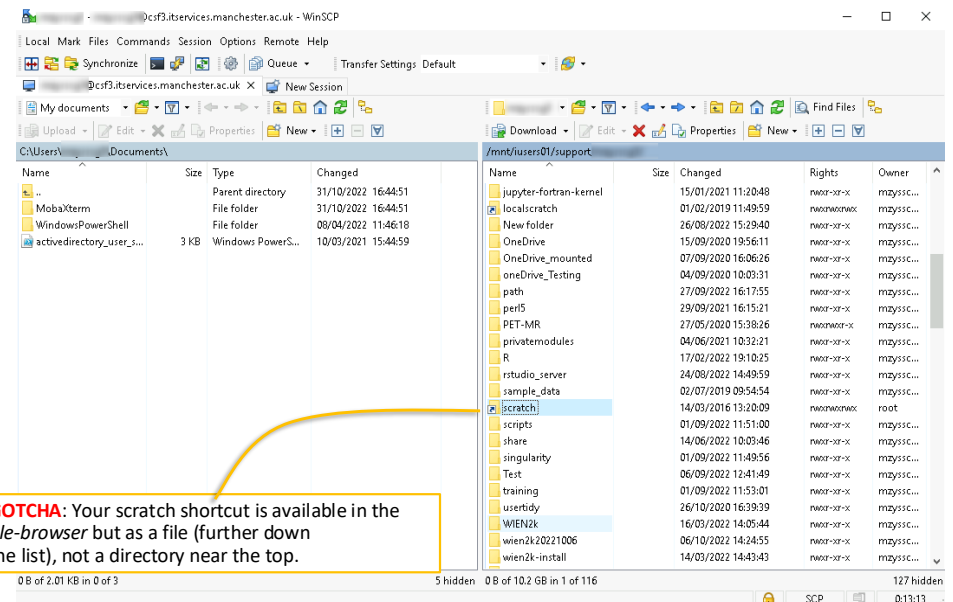
# Accessing scratch with MobaXterm



This can be flaky - do not use

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# Accessing scratch with WinSCP



GOTCHA: Your scratch shortcut is available in the file-browser but as a file (further down the list), not a directory near the top.

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# scratch file transfer with Linux / Mac

(can also type in a local MobaXterm window)

- Very similar to commands used earlier for our *home* directory

- Transfer a file from your computer to your CSF scratch dir

```
scp file2.txt username@csf3.itservices.manchester.ac.uk:~/scratch/
```

We give a destination after the : meaning "use my scratch shortcut"  
Omit the destination to transfer to *home* dir

- Transfer a file from your CSF scratch to your computer

```
scp username@csf3.itservices.manchester.ac.uk:scratch/results2.out results2.copy
```

Now you get a copy with a different name on your computer.  
Use . to keep the same name (results2.out)

- Change directory & filenames as required...

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## Extra Storage Space (Optional)

- Some research groups have extra space, example path:

```
/mnt/eps01-rds/group/username
```

- No shortcut from your home? To access it use:

```
cd /mnt/eps01-rds/group/username
```

- To create a shortcut (named *data*) in your home area:

```
cd ~
```

```
ln -s /mnt/eps01-rds/group/username data
```

- Also backed up
- Often many TB, but again shared by everyone else from your group
  - Be fair in your usage

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## Additional filesystem/file transfer info

- We have additional info about how to manage your files and your disk usage:  
<https://ri.itservices.manchester.ac.uk/userdocs/file-management/>
- Docs about file transfer:  
<https://ri.itservices.manchester.ac.uk/userdocs/file-transfer/>
- If you need to transfer a lot of files or big files to and from the CSF please do not do it on the login node
  - Ask for an account on the **RDS-SSH service**
- Many file management tasks can be included in your batch jobs – see the FAQ.

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## Basic Linux File Commands

A good Linux tutorial is available at: <https://www.chm.bris.ac.uk/unix/>

Command	Description
<pre>less file1 zless file2.gz</pre>	Display the content of <i>file1</i> (text file) a page at a time on screen. If you've compressed <i>file2</i> with <i>gzip</i> , no need to uncompress first. Press <i>space</i> to page down through a long file Press <i>return</i> to scroll down a line at a time Press <i>b</i> to scroll back up a page Press <i>G</i> to go to end of file Press <i>q</i> to quit/exit
<pre>cat file1 zcat file2.gz</pre>	Dump entire file to screen (a quick way to look at text files). If you've compress <i>file2</i> with <i>gzip</i> , no need to uncompress first.
<pre>gedit file1</pre>	Edit <i>file1</i> using a simple graphical text editor (similar to notepad on Windows). See later for more on opening graphical programs on the CSF so that they display a window on your computer.
<pre>file filenameA</pre>	Try to tell us what type of data is in <i>filenameA</i> . Useful to determine the output of some program where you are not sure what type of output it has generated. For example: <pre>file output.dat</pre> Might be ASCII text (so we can look at it with <i>less</i> or <i>gedit</i> ) or might be data (you'll need some other program to read it)
<pre>du -sh .</pre>	How much disk space is current directory (all files and subdirs) using?
<pre>df -h .</pre>	How much free space is there in the current area?

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## Basic Linux File Commands

A good Linux tutorial is available at: <https://www.chm.bris.ac.uk/unix/>

Command	Description
cd dir1 cd ~/dir1/dir2 cd .. cd	Change directory (go in to dir1 which is located in the current dir) Go in to dir2 in dir1 in home (~ is shorthand for <i>home</i> ) Go up to parent directory (e.g., from ~/dir1/dir2 to ~/dir1) Go back to <i>home</i> (useful if you become lost)
pwd	Lost? Print Working Directory (display current location)
ls ls -lh ls -lh file1 dirA ls -lh dirA/*.dat	List content (names of files and directories) of current directory List in long form (dates, file sizes, names) current directory List in long form (dates, file sizes, names) specified files, directories ... List in long form all files ending in .dat in directory dirA
mkdir dirA	Make directory named dirA (in the current directory)
cp fileA fileB	Copy (duplicate) a file (copy fileA to a new file fileB)
mv fileC fileD mv fileE dirA mv fileF dirA/fileG	Rename a file (from fileC to fileD). Works for directories too. Move fileE in to sub-directory dirA (dirA must exist) Move fileF AND rename it all in one go (dirA must exist)
rm fileH	Delete (remove) a file (caution!!)
rm -rf dir1	Delete directory and all files (and other sub-dirs) in there (caution!!!!)
gzip bigfile gunzip bigfile.gz	Compress a file (becomes bigfile.gz) to make better use of disk-space. Text files usually compress well. Uncompress previously compressed file (becomes bigfile).

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## PRACTICAL SESSION 3

File Transfer

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## Practical Session 3 – File Transfer

- Follow the hand-out 'Practical Session 3'
  - Transfer a file: from the CSF to your PC
  - Transfer a file: from your PC to the CSF
  - Windows: use MobaXterm, Mac/Linux: use "scp"
    - Or, if time permits, Windows users can try option 2 - <https://winscp.net/eng/download.php>
- This is not a 'real' world example, but:
  - You may need to generate files on your PC for processing on the CSF (e.g. an "abaqus" input file or a )
  - Your supervisor may give you files that you then need to transfer to CSF
- Exercise sheet (pdf) available at: <https://ri.itservices.manchester.ac.uk/course/rcsf/>

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## Need more help with the CSF?

- Extensive documentation about all aspects of the service:  
<https://ri.itservices.manchester.ac.uk/csf3/>
- Contact the Research Infrastructure Team  
[its-ri-team@manchester.ac.uk](mailto:its-ri-team@manchester.ac.uk)
- See you:
  - After lunch (in-person courses)

Thank you!

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